

SEQUENCE LISTING

<110> NAGOYA INDUSTRIAL SCIENCE RESEARCH INSTITUTE
GIFU INTERNATIONAL INSTITUTE OF BIOTECHNOLOGY
YAMADA, Yoshiji
YOKOTA, Mitsuhiro

<120> Method for diagnosing a risk of restenosis after percutaneous coronary intervention

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<150> JP P2002-233041

<151> 2002-08-09

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<210> 15
 <211> 1337
 <212> DNA
 <213> Homo sapiens

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<210> 16
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 <212> DNA
 <213> Artificial Sequence

<220>
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<220>
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 <222> (16)..(16)
 <223> n stands for any base

<400> 16
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<210> 17
 <211> 19
 <212> DNA
 <213> Artificial Sequence

<220>
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<220>
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 <223> n stands for any base

<400> 17
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<210> 18
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<220>
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<400> 18
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<210> 19
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<212> DNA
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24

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<400> 23
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<210> 24
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<220>
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<400> 24
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<400> 25
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18

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<212> DNA
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<220>
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<210> 38

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18

<210> 39
<211> 20
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<220>
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<400> 39
gtcacagtcg gtgccaatgt

20

<210> 40
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<212> DNA
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<220>
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<400> 40
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21

<210> 41
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<223> n stands for any base

<400> 41
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21

<210> 42
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<400> 42
ctgcagggaa gggagctgt

19

<210> 43
<211> 21
<212> DNA
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<400> 43
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21

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<223> n stands for any base

<400> 44
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22

<210> 45
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<212> DNA
<213> Artificial Sequence

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<400> 45

tcttacctga atctctgatc ttca

24

<210> 46
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<223> n stands for any base

<400> 46
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19

<210> 47
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<212> DNA
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<223> n stands for any base

<400> 47
cattcaccgt ggccangg

18

<210> 48
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<212> DNA
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<220>
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<400> 48
agctgcctgt accaatacat cc

22

<210> 49
<211> 22
<212> DNA
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<400> 49
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22

<210> 50
<211> 24
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<220>
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<223> n stands for any base

<400> 50
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24

<210> 51
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<212> DNA
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<220>
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<400> 51
caaaaacaac ttcaatgttt cga

23

<210> 52
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<212> DNA
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<400> 52
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16

<210> 53
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<210> 54
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<212> DNA
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<220>
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<400> 54
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<210> 55
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<212> DNA
<213> Artificial Sequence

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<400> 55
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<210> 56
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:Primer

<400> 56
ggccgcctcc gatgataca 19

<210> 57
<211> 24
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<400> 57
acccaaatac atctcccagg ancg

24

<210> 58
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<212> DNA
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<220>
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<223> n stands for any base

<400> 58
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24

<210> 59
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<212> DNA
<213> Artificial Sequence

<220>
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<400> 59
gaatgatatt gttgctgtgg gac

23

<210> 60
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<220>
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<223> n stands for any base

<400> 60
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19

<210> 61
<211> 19
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<220>
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<220>
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<400> 61
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19

<210> 62
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<212> DNA
<213> Artificial Sequence

<220>
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<220>
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<222> (12)..(12)
<223> n stands for any base

<400> 62
cacctgggcc antgcaggat

20

<210> 63
<211> 20
<212> DNA
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<220>
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<400> 63
cacctgggcc anggcaggat

20

<210> 64
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<220>
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<222> (9)..(9)
<223> n stands for any base

<400> 64
gaatcaagng cttttcgaaa catt

24

<210> 65
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
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<220>
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<222> (9)..(9)
<223> n stands for any base

<400> 65
gaatcaagna cttttcgaaa catt

24

<210> 66
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:Probe

<400> 66
tggacacgtg ggggagtcag

20

<210> 67
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:Probe

<400> 67
tggacacgtg gggagtcagc

20